<110> Stomp, Anne-Marie

SEQUENCE LISTING

```
Dickey, Lynn
      Gasdaska, John
<120> Expression of Biologically Active
  Polypeptides in Duckweed
<130> 40989/237225
<150> US 60/293,330
<151> 2001-05-23
<150> US 60/221,705
<151> 2000-07-31
<160> 8
<170> FastSEQ for Windows Version 4.0
<210> 1
<211> 554
<212> DNA
<213> Zea mays
<400> 1
gatcaagtgc aaaggtccgc cttgtttctc ctctgtctct tgatctgact aatcttggtt 60
tatgattcgt tgagtaattt tggggaaagc ttcgtccaca gttttttttt cgatgaacag 120
tgccgcagtg gcgctgatct tgtatgctat cctgcaatcg tggtgaactt atgtctttta 180
tatccttcac taccatgaaa agactagtaa tctttctcga tgtaacatcg tccagcactg 240
ctattaccgt gtggtccatc cgacagtctg gctgaacaca tcatacgata ttgagcaaag 300
atctatcttc cctgttcttt aatgaaagac gtcattttca tcagtatgat ctaagaatgt 360
tgcaacttgc aaggaggcgt ttctttcttt gaatttaact aactcgttga gtggccctgt 420
ttctcggacg taaggccttt gctgctccac acatgtccat tcgaatttta ccgtgtttag 480
caagggcgaa aagtttgcat cttgatgatt tagcttgact atgcgattgc tttcctggac 540
                                                                    554
ccgtgcagct gcgg
<210> 2
<211> 498
<212> DNA
<213> Artificial Sequence
<223> Duckweed codon optimized nucleotide sequence
      encoding human alpha-2B interferon
<221> CDS
<222> (1)...(498)
<400> 2
tgc gac ctc ccc cag acc cac agc ctc ggg tcc cgc cgc acc ctc atg
Cys Asp Leu Pro Gln Thr His Ser Leu Gly Ser Arg Arg Thr Leu Met
                                      10
                                                          15
```

ctg Leu	ctg Leu	gcg Ala	cag Gln 20	atg Met	cgc Arg	cgc Arg	atc Ile	tcg Ser 25	ctc Leu	ttc Phe	agc Ser	tgc Cys	ctg Leu 30	aag Lys	gac Asp	96
cgc Arg	cac His	gac Asp 35	ttc Phe	ggc Gly	ttc Phe	ccg Pro	cag Gln 40	gag Glu	gag Glu	ttc Phe	ggc Gly	aac Asn 45	cag Gln	ttc Phe	cag Gln	144
aag Lys	gcc Ala 50	gag Glu	acg Thr	atc Ile	ccc Pro	gtg Val 55	ctc Leu	cac His	gag Glu	atg Met	atc Ile 60	cag Gln	cag Gln	atc Ile	ttc Phe	192
aac Asn 65	ctg Leu	ttc Phe	agc Ser	acc Thr	aag Lys 70	gac Asp	agc Ser	tcg Ser	gcc Ala	gcc Ala 75	tgg Trp	gac Asp	gag Glu	acc Thr	ctg Leu 80	240
ctc Leu	gac Asp	aag Lys	ttc Phe	tac Tyr 85	acc Thr	gag Glu	ctg Leu	tac Tyr	cag Gln 90	cag Gln	ctc Leu	aac Asn	gac Asp	ctg Leu 95	gag Glu	288
gcg Ala	tgc Cys	gtg Val	atc Ile 100	cag Gln	ggg	gtt Val	ggg Gly	gtt Val 105	acg Thr	gag Glu	acg Thr	ccg Pro	ctg Leu 110	atg Met	aag Lys	336
gag Glu	gac Asp	agc Ser 115	atc Ile	ctc Leu	gcc Ala	gtg Val	cgc Arg 120	aag Lys	tac Tyr	ttc Phe	cag Gln	cgc Arg 125	ile	acg Thr	ctc Leu	384
tac Tyr	ctc Leu 130	aag Lys	gag Glu	aag Lys	aag Lys	tac Tyr 135	agc Ser	ccg Pro	tgc Cys	gcc Ala	tgg Trp 140	Glu	gtc Val	gtt Val	cgc Arg	432
gcc Ala 145	gag Glu	atc	atg Met	cgc Arg	tcc Ser 150	Phe	agc Ser	ctg Leu	agc Ser	acc Thr 155	Asn	cto Leu	cag Gln	gag Glu	agc Ser 160	480
				gag Glu 165	. *											498
<21 <21	<210> 3 <211> 96 <212> DNA <213> Oryza sativa															
acc	<400> 3 accatgcagg tcctgaacac gatggtcaac aagcacttcc tctccctgtc cgtcctcatc gtcctcctcg ggctgagcag caacctcacc gccggc															e 60 96
<21 <21	.0> 4 .1> 1 .2> E	L88 PRT	sapi	iens												
<4(Met	00> 4 : Ala	4 a Le	ı Thi	r Phe	e Ala	a Lei	ı Leı	ı Val	l Ala	a Le	u Lei	u Va	l Le	u Se:	r Cys	

```
10
Lys Ser Ser Cys Ser Val Gly Cys Asp Leu Pro Gln Thr His Ser Leu
          20
                               25
Gly Ser Arg Arg Thr Leu Met Leu Leu Ala Gln Met Arg Arg Ile Ser
                           40
Leu Phe Ser Cys Leu Lys Asp Arg His Asp Phe Gly Phe Pro Gln Glu
Glu Phe Gly Asn Gln Phe Gln Lys Ala Glu Thr Ile Pro Val Leu His
                   70
                                       75
Glu Met Ile Gln Gln Ile Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser
                                   90
Ala Ala Trp Asp Glu Thr Leu Leu Asp Lys Phe Tyr Thr Glu Leu Tyr
                              105
Gln Gln Leu Asn Asp Leu Glu Ala Cys Val Ile Gln Gly Val Gly Val
                          120
Thr Glu Thr Pro Leu Met Lys Glu Asp Ser Ile Leu Ala Val Arg Lys
                       135
Tyr Phe Gln Arg Ile Thr Leu Tyr Leu Lys Glu Lys Lys Tyr Ser Pro
                   150
                                       155
Cys Ala Trp Glu Val Val Arg Ala Glu Ile Met Arg Ser Phe Ser Leu
               165
                                   170
Ser Thr Asn Leu Gln Glu Ser Leu Arg Ser Lys Glu
```

<210> 5 <211> 165 <212> PRT <213> Homo sapiens

Cys Asp Leu Pro Gln Thr His Ser Leu Gly Ser Arg Arg Thr Leu Met 10 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp 25 Arg His Asp Phe Gly Phe Pro Gln Glu Glu Phe Gly Asn Gln Phe Gln 40 Lys Ala Glu Thr Ile Pro Val Leu His Glu Met Ile Gln Gln Ile Phe 55 Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr Leu 75 Leu Asp Lys Phe Tyr Thr Glu Leu Tyr Gln Gln Leu Asn Asp Leu Glu 85 90 Ala Cys Val Ile Gln Gly Val Gly Val Thr Glu Thr Pro Leu Met Lys 105 Glu Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr Leu 120 Tyr Leu Lys Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val Arg 135 140 Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu Ser

<210> 6 <211> 31

Leu Arg Ser Lys Glu

165

```
<212> PRT
<213> Oryza sativa
<400> 6
Met Gln Val Leu Asn Thr Met Val Asn Lys His Phe Leu Ser Leu Ser
                5
                                    10
Val Leu Ile Val Leu Leu Gly Leu Ser Ser Asn Leu Thr Ala Gly
<210> 7
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> Modified rice alpha-amylase signal peptide
<400> 7
Met Gln Val Leu Asn Thr Met Val Asn Lys His Phe Leu Ser Leu Ser
                5
                                    10
Val Leu Ile Val Leu Thr Val Leu Ser Ser Asn Leu Thr Ala Gly
                                25
<210> 8
<211> 21
<212> PRT
<213> Arabidopsis thaliana
<400> 8
Met Lys Thr Asn Leu Phe Leu Phe Leu Ile Phe Ser Leu Leu Ser
                                    10
Leu Ser Ser Ala Glu
            20
```